Manual **Analysis**

plateCore Workflow

Import raw FCS files

> pbmcFP <- flowPlate(pbmcPlate,</pre> wellAnnotation,plateName="PBMC.001")

Per well compensation

> pbmcFP <- compensate(pbmcFP,compensation.matrix)</pre>

Select population(s) of interest



> pbmcFP <- Subset(pbmcFP, rectangleGate(

"FSC-H"=c(300,700),"SSC-H"=c(50,400)))

Per plate quality control



> ecdfplot(~`FSC-H`|as.factor(Row.ld),plateSet(pbmcFP))

Set control gates



> pbmcFP <- setControlGates(pbmcFP,</p> gateType="Negative.Control")

Score test samples



> pbmcFP <- applyControlGates(pbmcFP)</p>

Annotation and analysis

> pbmcFP <- summaryStats(pbmcFP)</pre>